

**Fig. 1**

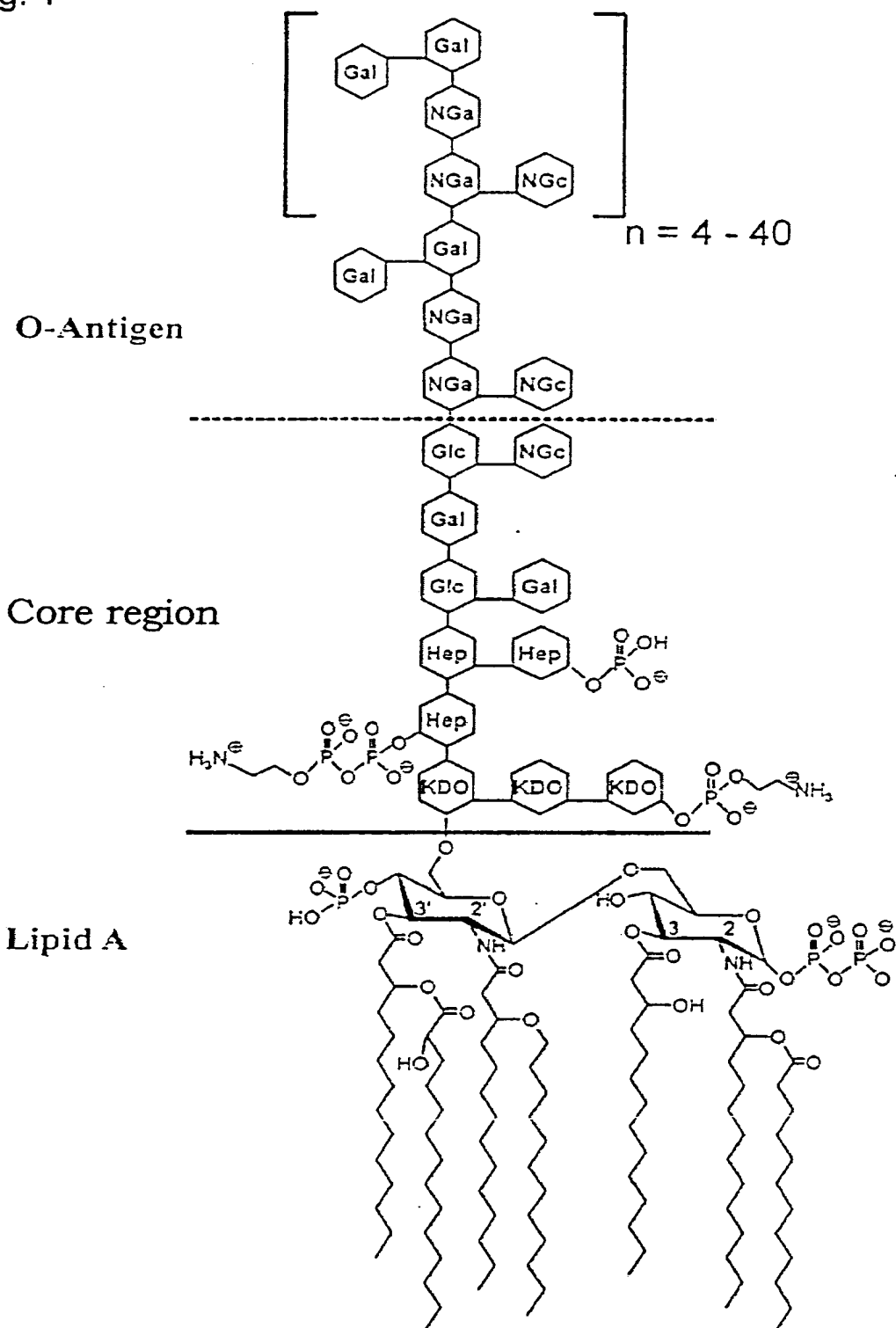


Fig. 2

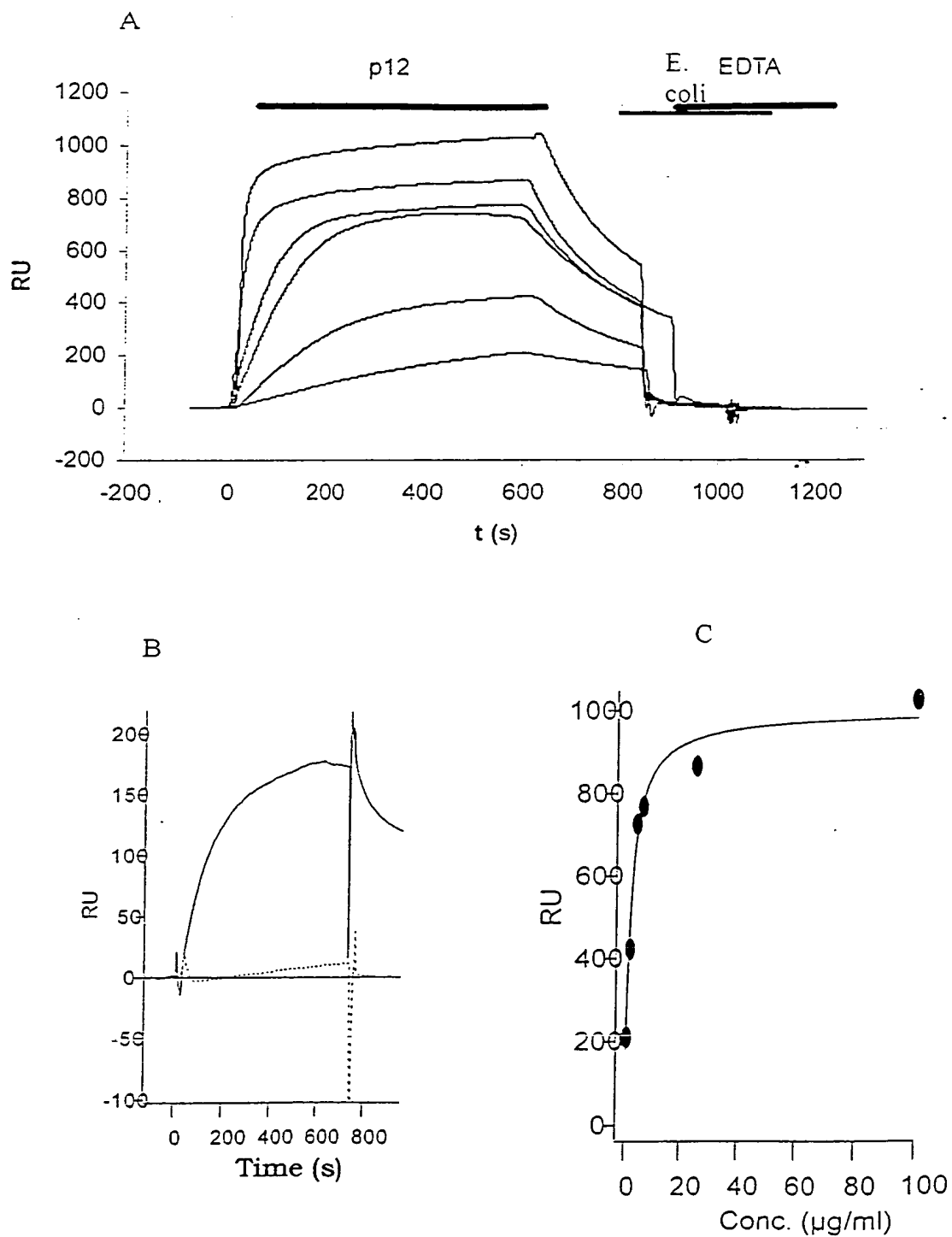
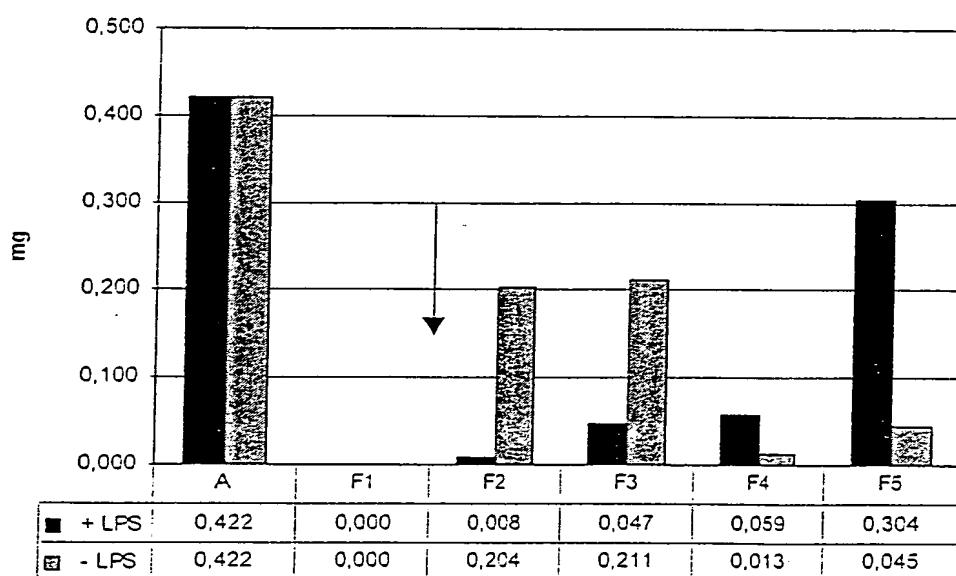


Fig. 3

Endotoxin structure	<i>E. coli</i> strain	p-12 bond
<pre>       KDO-LipidA               KDO               KDO           </pre>	D21f2	-
<pre>       Hep-Hep-KDO-LipidA                   Hep  KDO                       KDO           </pre>	D21f1	+
<pre>       Glc-Hep-Hep-KDO-LipidA                   Hep  KDO                       KDO           </pre>	D21e8	+
<pre>       Glc-Hep-Hep-KDO-LipidA                   Gal Hep  KDO                       KDO           </pre>	D21e7	+
<pre>       GlcN-Glc-Glc-Glc-Glc-Hep-Hep-KDO-LipidA                   Gal Hep  KDO                       KDO           </pre>	D21	+

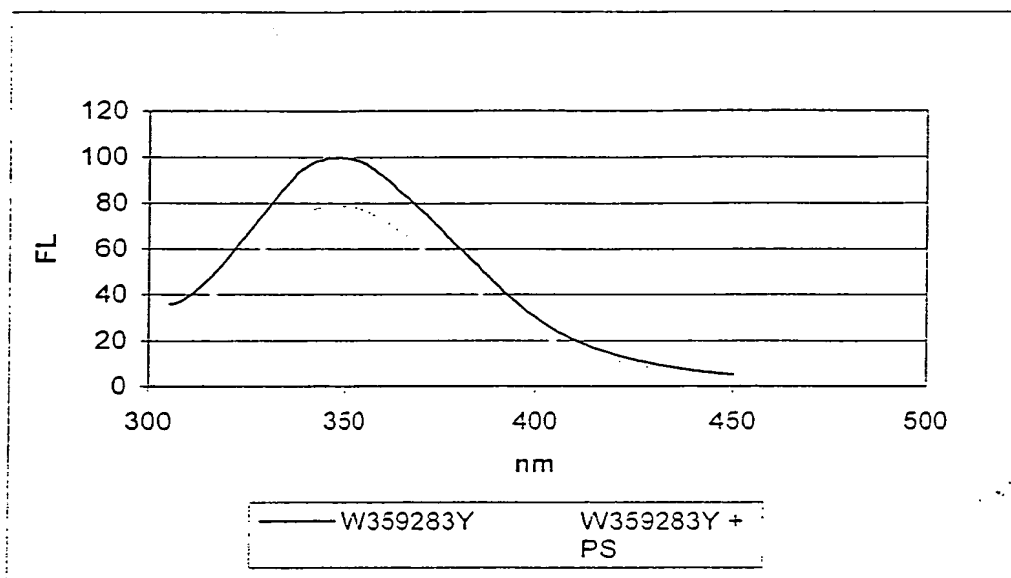
pH	K <sub>d</sub>
6,0	3,09 E-07
7,5	6,85 E-08
8,0	5,86 E-08
8,5	7,86 E-08
9,0	3,29 E-08
10,0	1,55 E-07

Fig. 4

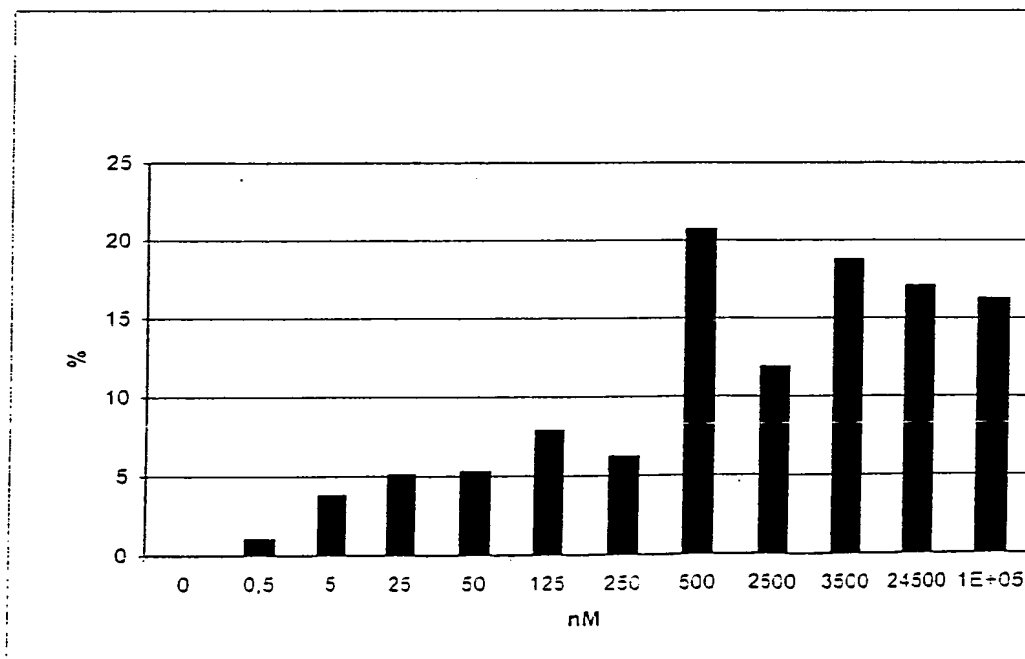


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Fig. 5 A

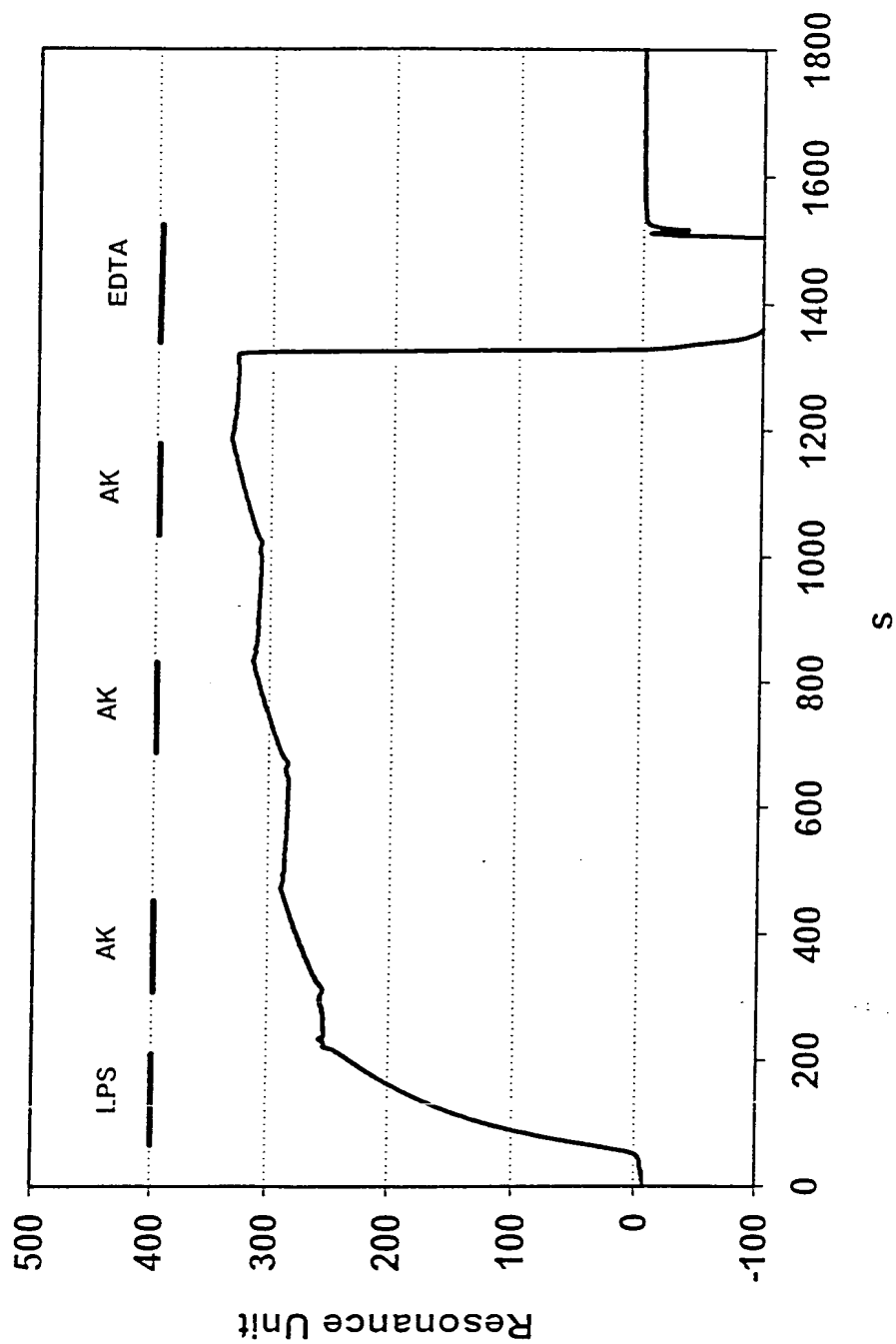


B



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Fig. 6



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Fig. 7A

T2p12	MSNNTYQHVS NESRYVKFDPTDTNFPPEITDVQAAIAAISPAGVNGVPDASSTTKGILFL
K3p12	MSNNTYQHVS NESRYVKFDPTDTNFPPEITDVQAAIAAISPAGVNGVPDASSTTKGILFL
T4p12	MSNNTYQHVS NESRYVKFDPTDTNFPPEITDVHAAIAAISPAGVNGVPDASSTTKGILFI
RB32-33p12	MSNNTYQHVS NESKYVKFDPVGSNFPDPTVTTVQSALS KISNIGVNGIPDATMEVKGIAMI
AR1p12	MSNNTYQHVS NESKYVKFDPVGSNFPDPTVTTVQSALS KISNIGVNGIPDATMEVKGIAMI
PP01p12	MSNNTYQHVS NESKYVKFDPVGSNFPDPTVTTVQSALS KISNIGVNGIPDASMEVKGIAMI
RB69p12	MSNNTYQHVS NESVYVEFDPTGSNFDSSITNVQAALASISAYGVKGVPDASEAEKGV IQL ***** **:
T2p12	ATEQEVIDGTNNTKAVTPATLATRLSYPNATEAVYGLTRYSTDDEAIAGVNNESSITPAK
K3p12	ATEQEVIDGTNNTKAVTPATLATRLSYPNATETVYGLTRYSTNDEAIAGVNNESSITPAK
T4p12	PTEQEVIDGTNNTKAVTPATLATRLSYPNATETVYGLTRYSTNDEAIAGVNNESSITPAK
RB32-33p12	ASEQEVL DGTNNSKI VTPATLATRLLYPNATETKYGLTRYSTNEETLEGS DNNSSITPQK
AR1p12	ASEQEVL DGTNNSKI VTPATLATRLLYPNATETKYGLTRYSTNEETLEGS DNNSSITPQK
PP01p12	ASEQEVL DGTNNSKI VTPATLATRLLYPNATETKYGLTRYSTNEETLEGS DNNSSITPQK
RB69p12	ATEQEVL DGFNSTKAVTPATLNARLQYPNASETQYGVTKYATQEEAIAGTLDTVSITPLK .:****:
T2p12	FTVALNNVFETR VSTESSNGVIKISSLPQALAGADDTTAMTPLKTQQLAVKLI AQIAPSK
K3p12	FTVALNNAFETR VSTESSNGVIKISSLPQALAGADDTTAMTPLKTQQLAIKLI AQIAPSE
T4p12	FTVALNNAFETR VSTESSNGVIKISSLPQALAGADDTTAMTPLKTQQLAIKLI AQIAPSE
RB32-33p12	LKYHTDDVFQNRYSSESSNGVIKISSTPAALAGVDDTTAMTPLKTQKLAIKLISQIAPSE
AR1p12	LKYHTDDVFQNRYSSESSNGVIKISSTPAALAGVDDTTAMTPLKTQKLAIKLISQIAPSE
PP01p12	LKYHTDDVFQNRYSSESSNGVIKISSTPAALAGVDDTTAMTPLKTQKLAIKLISQIAPSE
RB69p12	LNQ TIDNTFSTRYSTETTNGVIKIATQTAALAGSDDTTAMTPLKTQQLAIKLISQIAPNN :..*:
T2p12	NAATESEQV IQLATVAQARQGT LREGY AISP YTFMNSTATEEYKGV IKLGTQSEVNSNN
K3p12	TTATESDQGVVQLATVAQVRQGT LREGY AISP YTFMNSSATEEYKGV IKLGTQSEVNSNN
T4p12	TTATESDQGVVQLATVAQVRQGT LREGY AISP YTFMNSSATEEYKGV IKLGTQSEVNSNN
RB32-33p12	DTASESVRGVVQLSTVAQTRQGT LREGY AISP YTFMNSVATQEYKGV IRLGTQSEINSNL
AR1p12	DTASESVRGVVQLSTVAQTRQGT LREGY AISP YTFMNSVATQEYKGV IRLGTQSEINSNL
PP01p12	DTASESVRGVVQLSTVAQTRQGT LREGY AISP YTFMNSVATQEYKGV IRLGTQSEINSNL
RB69p12	DPASEITGVVRLATVAQTRQGT LREGY AISP YTFMNSVATQEYKGV IRLGTQAEINSNL .*:
T2p12	ASVAVTGATLNGRGSTTSMRGVVKLTTTAGSQSGGDASSALAWNADVIHQ RGGQTINGTL
K3p12	ASVAVTGATLNGRGSTTSMRGVVRLTTTAGSQSGGDASSALAWNADVIHQ RGGQTINGTL
T4p12	ASVAVTGATLNGRGSTTSMRGVVKLTTTAGSQSGGDASSALAWNADVIQQRGGQIIYGT L
RB32-33p12	GDVAVTGETLNGRGATSSMRGVVKLTTQAGIAPEGDGSGALAWNADVINTRGGQTINGSL
AR1p12	GDVAVTGCTLNGRGATGSMRCVVKLTTQAGIAPEGDSSGALAWNADVINTRGGQTINGSL
PP01p12	GDVAVTGETLNGRGATGSMRGVVKLTTQAGIAPEGDSSGALAWNADVINTRGGQTINGSL
RB69p12	GDVAVTGETLNGRGATGSMRGVVKLTTQAGVAPEGDSSGALAWNADVINTRGGQTINGSL ..***** *****:* *****:*:* *
T2p12	RINNTLTIASGGANITGTVNMTGGYIQGKR VVTQNEIDRTIPVGAIMMWAADSLPSDAWR
K3p12	RINNTLTIASGGANITGTVNMTGGYIQGKR VVTQNEIDRTIPVGAIMMWAADSLPSDAWR
T4p12	RIEDTFTIANGGANITGTVRMTGGYIQGNRI VVTQNEIDRTIPVGAIMMWAADSLPSDAWR
RB32-33p12	NLD---HLTANGIWSRGGMWKNG---DQPVATERYASERVPVGTIMMFAGDSAP-PGWI
AR1p12	NLD---HLTANGIWSRGGMWKNG---DQPVATERYASERVPVGTIMMFAGDSAP-PGWI
PP01p12	NLD---HLTANGIWSRGGMWKNG---DQPVATERYASERVPVGTIMMFAGDSAP-PGWI
RB69p12	NLD---HLTANGIWSRGGMWKNG---DQPVATERYASERVPVGTIQMFAGDSAP-PGVV .: : *

Fig. 7B

T2p12	FCHGGTVSASDCPLYASRIGTRYGGTSSNPGLPDMRGLFVRGSGRGSHTLNPVNNGNDQF
K3p12	FCHGGTVSASDCPLYASRIGTRYGGSSSNPGLPDMRGLFVRGSGRGSHTLNPVNNGNDQF
T4p12	FCHGGTVSASDCPLYASRIGTRYGGNPSNPGLPDMRGLFVRGSGRGSHTLNPVNNGNDQF
RB32-33p12	MCHGGTVSGDQYPDYRNTVGARFGGDWNNPGIPDMRGLFVRGAGTGGHILNQ--RGQDGY
AR1p12	MCHGGTVSGDQYPDYRNTVGTRFGGDWNNPGIPDMRGLFVRGAGTGGHILNQ--RGQDGY
PP01p12	MCHGGTVSGDQYPDYRNTVGTRFGGDWNNPGIPDMRGLFVRGAGTGXHILNQ--RGQDGY
RB69p12	LCHGGTISGDQFPDYRNVVCTRFGGDWNPNPGIPDMRGLFVRGAGTGSHILNN--RGQDGY
	:*****:*.:.: * * . :*:**:** .***:*****:*. * *: * .*: * :
T2p12	GKPRLGVGCTGGYVGEVQKQQMSYHKHAGGFGEY---DDSGAFGNTRRSNFGVTRKGLDW
K3p12	GKPRLGVGCTGGYVGEVQKQQMSYHKHAGGFGEW---DDSGAFGNTRRSNFGVTRKGLDW
T4p12	GKPRLGVGCTGGYVGEVQIQQMSYHKHAGGFGEH---DDLGAFGNTRRSNFGVTRKGLDW
RB32-33p12	GKDRLGVGCDGMHVGGVQAQQMSYHKHAGGWGEY--QRHEAPFGASVYQGYLGTRKYSW
AR1p12	GKDRLGVGCDGMHVGGVQAQQMSYHKHAGGWGEY--NRSEGPFGASVYQGYLGTRKYSW
PP01p12	GKDRLGVGCDGMHVGGVQAQQISYHKHAGAWGENGNRGYAPFGASNGSGYLGNGRSADW
RB69p12	GKDRLGVGCDGMHVGGVQAQQMSYHKHAGGWGEF--QRHEAPFGASVYQGYLGTRKYSW
	** ***** * :** ** **:*****:.* ..** : ..:*. : **
T2p12	DNRSYFTNDGYEIDPASQRNSRYTLNRPELIGNETRPWNISLNYIIKVKE
K3p12	DNRSYFTNDGYEIDPASQRNSRYTLNRPELIGNETRPWNISLNYIIKVKE
T4p12	DNRSYFTNDGYEIDPESQRNSKYTLNRPELIGNETRPWNISLNYIIKVKE
RB32-33p12	DNASYFTNDGFELG--GPRDALGTLNREGLIGYETRPWNISLNYIIKIHY
AR1p12	DNASYFTNDGFELG--GPRDALGTLNREGLIGYETRPWNISLNYIIKIHY
PP01p12	DNHLFFTNDGFEMG--GPRDSFGTLNREGLIGYETRPWNISLNYIIKIHY
RB69p12	DNASYFTNDGFELG--GHRDATGTLNREGLIGYETRPWNISLNYIIKVHY
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